

Dual Role of Oxidized LDL on the NF-KappaB Signaling Pathway

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Atherosclerosis is a slowly evolutive age-linked disease of large arteries, characterized by a local lipid deposition associated with a chronic inflammatory response, leading potentially to acute plaque rupture, thrombosis and ischemic heart disease. Atherogenesis is a complex sequence of events associating first expression of adhesion molecules, recruitment of mononuclear cells to the endothelium, local activation of leukocytes and inflammation, lipid accumulation and foam cell formation. Low density lipoproteins (LDLs) become atherogenic after undergoing oxidation by vascular cells, that transform them into highly bioreactive oxidized LDL (oxidized LDLs). Oxidized LDLs are involved in foam cell formation, and trigger proatherogenic events such as overexpression of adhesion molecules, chemoattractant agents growth factors and cytokines involved in the inflammatory process, cell proliferation and apoptosis. Moreover, this toxic effect of oxidized LDLs plays probably a role in plaque erosion/rupture and subsequent atherothrombosis.

Several biological effects of oxidized LDLs are mediated through changes in the activity of transcription factors and subsequently in gene expression. Oxidized LDLs exert a biphasic effect on the redox-sensitive transcription factor NF- κ B, which can be activated thereby up-regulating proinflammatory gene expression, such as adhesion molecules, tissue factor, scavenger receptor LOX-1. On the other hand, higher concentrations of oxidized LDLs may inhibit NF- κ B activation triggered by inflammatory agents such as LPS, and may thereby exert an immunosuppressive effect. This review is an attempt to clarify the mechanism by which oxidized LDLs may up- or down-regulate NF- κ B, the role of NF- κ B activation (or inhibition), and the consequences of the oxidized LDLs-mediated NF- κ B dysregulation and their potential involvement in atherosclerosis.

Keywords: Oxidized LDL; NF- κ B; Inflammation; Atherosclerosis; Reactive oxygen species

Abbreviations: Oxidized LDLs, oxidized low density lipoproteins; ROS, reactive oxygen species; NF- κ B, nuclear factor kappaB;

I κ B, IkappaB; VCAM-1, vascular cell adhesion molecule-1; ICAM-1, intracellular cell adhesion molecule-1; PPARs, peroxisome-proliferator-activated receptors

INTRODUCTION

Atherosclerosis with its related complications, heart attack, stroke and peripheral vascular diseases, is a prevalent cause of morbidity and mortality in western countries. Atherogenesis is characterized by cholesterol esters accumulation in atherosclerotic plaques, associated with several cellular effects, such as endothelial activation, migration of mononuclear cells, local chronic inflammatory process, smooth muscle cell proliferation and cell death. This may lead to acute complications, such as plaque erosion or rupture and thrombotic events.^[1–3]

Low density lipoproteins (LDLs) are involved in the pathogenesis of atherosclerosis after undergoing oxidative modifications in the arterial wall.^[3,4] Oxidized LDLs exhibit *in vitro* a variety of biological properties potentially involved in atherogenesis, such as lipid accumulation in macrophagic cells (foam cell formation), overexpression of adhesion proteins (recruitment of mononuclear cells), local immune response alteration (inflammatory response), smooth muscle cell migration and proliferation, production of extracellular matrix, alterations of coagulation pathways, disturbance in the arterial tone regulation, cytotoxicity (apoptosis or necrosis).^[4,5] These biological responses triggered by oxidized LDLs are mediated by various intracellular signaling pathways, and (dys-)regulation of gene

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expression.^[5-7] This led Hajjar and Haberland^[6] to consider oxidized LDLs as “trojan horses” because bioactive lipids internalized inside the cell trigger (dys-)regulation of various intracellular signaling pathways and of gene expression.^[5,6]

The inflammatory response is thought to play a major role in the atherogenesis and in the trigger of plaque erosion or rupture. NF- κ B is a central player in the regulation of both inflammatory response and cell survival. It is activated by a variety of factors (e.g. cytokines, oxidants) and also by oxidized LDLs. In fact, oxidized LDLs trigger a biphasic effect on NF- κ B, depending on their concentration.^[6] NF- κ B is rapidly activated in vascular cultured cells by low concentrations of oxidized LDLs, under conditions inducing mitogenic and proinflammatory responses.^[8] In contrast, longer periods of incubation and high oxidized LDLs concentrations impair NF- κ B activation mediated by various agonists, and may thereby induce proapoptotic and immunosuppressive effects.^[9] So far, how oxidized LDLs modulate cell signaling and mechanisms of NF- κ B activation, is only partly understood.

This review is an attempt to give a synthetic view of the effects of oxidized LDLs on the NF- κ B system, the mechanisms of activation and inhibition and the potential involvement in atherosclerosis.

THE NF- κ B SYSTEM

The transcription factor NF- κ B is an ubiquitous transcription factor that can be activated by a large number of extracellular stimuli such as cytokines, chemokines, growth factors, bacterial or viral products, hypoxia-reperfusion and stress generating responses.^[10-12] NF- κ B activation triggers the induction of inflammatory genes and may thereby play a role in the initiation and progression of various chronic inflammatory diseases and atherosclerosis.^[9,10]

The NF- κ B Family

The Rel/NF- κ B family consists of the members NF- κ B-1 (p50/p105), NF- κ B-2 (p52/p100), p65 (RelA), RelB and c-Rel which form various homodimers and heterodimers, where the most common active form is p50 or p52/p65.^[10-12] A characteristic feature of NF- κ B proteins is that all the family members share a highly conserved Rel homology domain (RHD). This domain, composed of approximately 300 amino acid residues, contains a nuclear localization sequence (NLS) and is responsible for dimerization and DNA binding. In addition, the RHD is the site for the binding of the NF- κ B inhibitors, I κ Bs.

Mechanisms of NF- κ B Activation

The I κ B family is constituted of (at least) seven members that share multiple copies of the ankyrin

motif. I κ B α , I κ B β , I κ B ϵ , I κ B γ /p100, I κ B δ /p105 and Bcl-3 control the activity of NF- κ B. In resting cells, the ankyrin repeats of I κ Bs mask the NLS of NF- κ B and prevent its nuclear translocation. Upon activation, I κ B is rapidly phosphorylated on two serine residues (ser-32 and ser-36 in I κ B α , ser-19 and ser-23 in I κ B β). This leads to I κ B ubiquitination and subsequent degradation by the 26S proteasome. The released NF- κ B dimer then translocates to the nucleus where it binds to specific promoters and activates the transcription of a wide variety of target genes (Fig. 1), including genes of cytokines (IL-1, TNF- α), adhesion molecules (VCAM-1, ICAM-1, E-selectin), regulators of apoptosis (c-IAPs, Bcl-xL), immunoreceptors (Table I).^[13,14]

The critical regulatory step in the activation of NF- κ B is the phosphorylation of the N-terminal regulatory serines of I κ B proteins which is mediated by a high molecular weight I κ B kinase complex (IKK) (approximately 700 kDa).^[15-17] IKK is a serine-specific kinase composed of at least three subunits: two catalytic subunits IKK α (IKK-1) and IKK β (IKK-2) and a regulatory subunit IKK γ . The predominant form of IKK is an heterodimer IKK α /IKK β associated with either dimer or trimer of IKK γ . A fourth protein, IKAP (IKK complex-associated protein) may also be involved in IKK activation, but its physiological function is not clearly understood.

The inactive (non phosphorylated) IKK complex is activated by phosphorylation mediated by an heterogeneous group of serine/threonine kinases, named “IKK-kinases”. This group includes members of the mitogen-activated protein kinase kinase kinase (MAPKKK) family such as NIK (NF- κ B inducing kinase) and MEKK-1, -2 and -3 (MAPK/ERK kinase kinase),^[17] protein kinase C (PKC),^[18] AKT/protein kinase B^[19] and two related I κ B-kinase IKKi/IKK ϵ and NAK/TBK1 (NF- κ B activating kinase/Tank binding kinase 1)^[20] (Fig. 1).

In response to upstream stimuli, IKK-kinases are activated and recruited to the complex via IKK γ . Activated IKK-kinases induce both phosphorylation of the catalytic subunits and activation of the IKK complex, which in turn phosphorylates I κ B. IKK activation is transient and is regulated by a C-terminal autophosphorylation and inactivation by phosphatases.^[15,16] A second mechanism of autoregulation of the NF- κ B/I κ B system is also induced by NF- κ B itself which elicits the synthesis of its own inhibitor I κ B. The newly synthesized I κ B α proteins enter the nucleus to bind activated NF- κ B and carry it to the cytoplasm, thereby terminating the activation of gene expression.

It may be noted that NF- κ B can be activated independently of the IKK complex by two alternative mechanisms. The first one is UV radiation-induced NF- κ B activation to which does not involve I κ B α phosphorylation. The second exception is anoxia, which stimulates phosphorylation of I κ B α

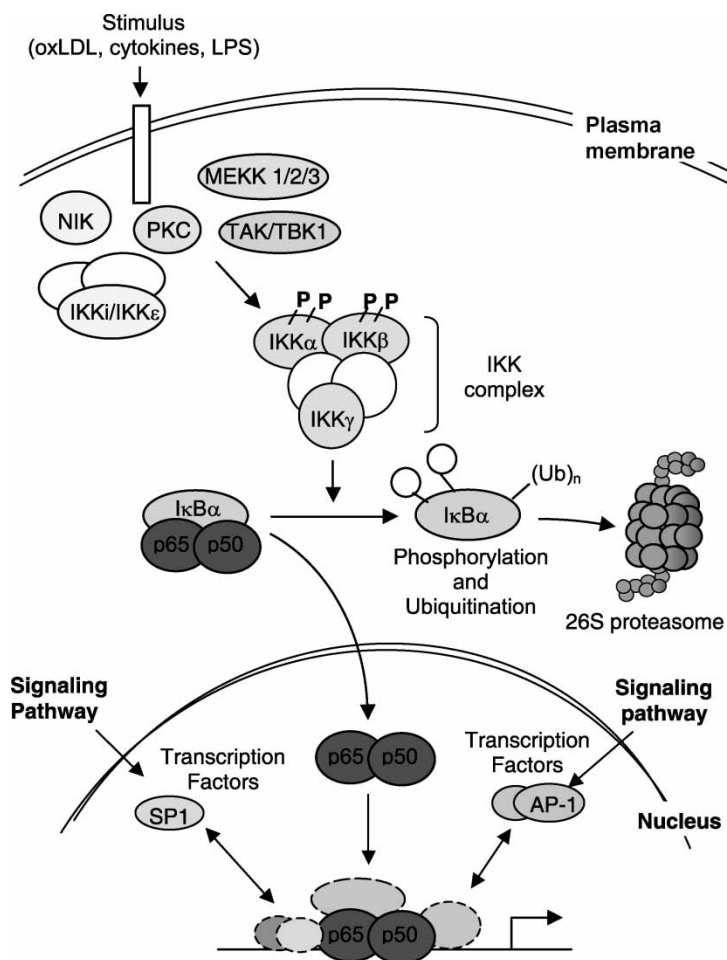


FIGURE 1 NF-κB signaling pathway.

at tyrosine 42, that may bind the SH2 domain of PI3-kinase which pulls it away from NF-κB.^[15-17]

NF-κB Activation by Oxidative Stress

NF-κB is a redox-sensitive transcription factor which is activated by intracellular reactive oxygen species

(ROS), such as H₂O₂, superoxide anion, hydroxyl radicals. This hypothesis is supported by the activating effect of exogenous H₂O₂ in various cell types and by the inhibitory effect of antioxidants, including both chemical antioxidants (such as NAC or PDTC) and overexpressed antioxidant enzymes.^[21-24]

TABLE I NF-κB target genes

Proinflammatory cytokines	IL-1 α-β, IL-2, IL-6, IL-12, TNF-α, LTα, LTβ, IFNβ, GM-CSF, M-CSF (Macrophage colony stimulating Factor), G-CSF (Granulocyte colony stimulating Factor)
Chemokines	IL-8, MiP-1 α (Macrophage inflammatory protein 1 α), MCP-1 (Macrophage chemotactic protein 1), RANTES, Eotaxine, Gro α, β, γ
Adhesion molecules	ICAM-I, VCAM-1, E-selectin
Acute phase proteins	SAA (Serum Amyloid A)
Inflammatory enzymes	iNOS (inducible Nitric Oxyde Synthase), COX-2 (cyclooxygenase-2), 5-lipoxygenase, cytosolic phospholipase A ₂
Receptors	Interleukin-2 receptor (α-chain), T-cell receptor (β-chain), Platelet-activating factor receptor (PAF-R)
Regulators of apoptosis	c-IAP-1, c-IAP-2, A1 (BFI-1), Bcl-X _L , Fas ligand, Fas (CD95), c-myc, p53, cycline D1, A20
Protein of coagulation	Tissue factor

See reviews: (Barnes, *New England J. of Medicine*, 1997—Ghost and Karin, *Cell*, 2002—May and Ghost, *Immunology Today*, 1998).

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The role played by ROS in the mechanism of NF- κ B activation remains only poorly understood.^[21] ROS could act as second messengers that modulate the kinase activities of NF- κ B signaling pathways, IKK complex activation^[22] and/or DNA binding and transactivation activity of the NF- κ B dimers. This role remains still cell and stimulus specific.^[23,24]

NF- κ B, OXIDIZED LDLs AND ATHEROSCLEROSIS

Several lines of evidence suggest that NF- κ B could contribute to the initiation and progression of atherosclerosis. Activated NF- κ B (which is not detected in the normal arterial wall), is present in areas predisposed to atherosclerotic lesion formation, in atherosclerotic lesions and in plaques of patients with unstable angina pectoris.^[25–28] The presence of activated NF- κ B in atherosclerotic areas raises the question of the potential role of NF- κ B in atherogenesis, since a local inflammatory process is implicated in atherogenesis and NF- κ B is known to be implicated in the inflammatory response.

Hypercholesterolemia and Oxidized LDLs are Associated with NF- κ B Activation

Several reports indicate that NF- κ B activation is associated with hyperlipidemia or oxidized LDLs. For instance, mice placed on an atherogenic diet or injected with oxidized LDLs exhibited an hepatic accumulation of lipid peroxides, associated with NF- κ B activation in the liver and the expression of a set of inflammatory and oxidative stress responsive genes, such as serum amyloid A (SAA), heme oxygenase and JE (the murine homologue of Macrophage Chemotactic protein-1, MCP-1).^[29,30] In coronary arteries of hypercholesterolemic pigs, activated NF- κ B colocalizes with areas of intense lipid staining.^[31] In a rat model, injected LDLs induced NF- κ B activation and expression of intracellular cell adhesion molecule-1 (ICAM-1) in aortic endothelium.^[32] It may be noted that a causal link between NF- κ B activation and early atherosclerotic lesions has not been firmly demonstrated so far, because genetically engineered mice deficient in p65 or I κ B α die during embryonic development.^[33]

Oxidized LDLs Components Involved in the Modulation of NF- κ B Activity

Various bioactive oxidized lipids present in oxidized LDLs are able to trigger NF- κ B activation.

13-HPODE (13-hydroperoxyoctadecadienoic acid) induces NF- κ B activation and subsequent overexpression of the vascular cell adhesion molecule-1 (VCAM-1) in porcine vascular SMCs.^[34] This effect is

mediated by oxidative stress and PKC as assessed by the inhibition of NF- κ B DNA binding and transcription of VCAM-1 gene by the antioxidant NAC and calphostin C.^[34]

Oxidized phospholipids isolated from minimally oxidized LDL induced monocyte binding on endothelium and increase in VCAM-1 expression.^[35,36]

Lysophosphatidylcholine (Lyso-PC) elicited PKC and tyrosine kinases activation and subsequent NF- κ B activation and overexpression of ICAM-1, VCAM-1 and MCP-1.^[37–40]

Oxysterols (7-keto, 7-hydroxy and 25-hydroxycholesterol are the major oxysterols present in oxidized LDLs) are thought to be responsible for a large part of the biological effects of oxidized LDLs on cultured cells, for instance, regulation of cholesterol metabolism, apoptosis, inflammation, immunosuppression and atherogenesis.^[41] To date NF- κ B activation by oxysterol has not been reported, but cannot be excluded *a priori*, since oxysterols induce a cellular oxidative stress (which is known to trigger NF- κ B activation) and trigger nuclear receptors LXR α and LXR β activation (which are often activated concomitantly with NF- κ B).^[42–44]

Hydroxynonenal (HNE) and other aldehydic products, formed during oxidation of polyunsaturated fatty acids (PUFA) of LDLs, are able to trigger or modulate cell signaling involved in proliferation or apoptosis.^[45] Low HNE concentration induces MAPK activation and AP-1 nuclear binding, but does not activate NF- κ B in these conditions.^[46–49] Moreover, HNE has been shown to downregulate NF- κ B nuclear binding and counteract lipopolysaccharide (LPS)-induced NF- κ B activation.^[49]

Mechanism of NF- κ B Activation Induced by Oxidized LDLs

Moderate concentrations of oxidized LDLs trigger a rapid and dose-dependent activation of NF- κ B.^[50] Oxidized LDLs trigger the activation of the PI-3K/Akt pathway which is involved in cell survival.^[51] The PI-3K/Akt pathway has been shown to associate with and activate the IKK complex^[52] and to increase the transactivation potential of NF- κ B (p65).^[53] However, in smooth muscle cells overexpressing mutated forms of PI-3K and AKT, oxLDLs are still able to induce NF- κ B activation, thus suggesting that other signaling pathways may be involved in the oxLDL-mediated NF- κ B nuclear translocation.^[46] This activation is concomitant with cell signaling leading to cell proliferation or death and may constitute a defense mechanism against apoptosis.

Another survival mechanism that regulates cell death and proliferation induced by oxidized LDLs

is the generation of bioactive sphingolipid metabolites, for instance sphingosine-1-phosphate (S1-P).^[54] S1-P is involved in NF- κ B activation mediated by inflammatory cytokines.^[55,56] The mechanism of S1-P-induced NF- κ B activation is poorly defined, but could involve calcium mobilization and oxidative stress.^[57] However, it may be noted that the oxidized LDL-induced S1-P generation in smooth muscle cells^[58] is apparently dispensable for NF- κ B activation, since S1-P inhibitors did not abrogate the oxidized LDL-induced NF- κ B activation.^[46]

ROS are also implicated in the oxidized LDL-induced NF- κ B activation, as assessed by the inhibitory effect of antioxidants.^[46,59,60] ROS are generated by a DPI-sensitive oxidase system activated through signaling triggered by binding of oxidized LDLs to LOX-1 (lectin-like ox-LDL receptor-1) in endothelial cells, since an anti-LOX-1 antibody inhibits significantly NF- κ B activation.^[59,60] How oxidized LDL-induced ROS activate NF- κ B is not completely elucidated, and various mechanisms may be involved. A first mechanism may implicate the PI-3K/Akt pathway which is activated by oxidized LDLs^[51] and is able to activate the IKK complex and NF- κ B (p65).^[52,53] Another mechanism may involve MAPK and the serine-threonine kinase RSK (pp90rsk). The serine-threonine kinase RSK (pp90rsk), a downstream target of MAPK, plays a role in the regulation of a number of cellular functions and cell survival via association and phosphorylation of signaling proteins, such as Bad and I κ B α , and of transcriptional regulators including c-Fos, estrogen receptor, NF- κ B, cAMP-response element-binding protein (CREB) and CREB-binding protein.^[61,62] As oxidized LDLs are able to activate MAPK,^[63] it may be hypothesized that they may also activate RSK (a well known downstream target of MAPK). However, RSK activation by oxidized LDLs has not been reported so far.

Finally, the ubiquitin/proteasome system is also required for I κ B degradation and NF- κ B nuclear translocation triggered by oxidized LDLs.^[46]

NF- κ B Inhibition by Oxidized LDLs

Unexpectedly, oxidized LDLs induce a biphasic effect on NF- κ B, since short term incubation of cells with oxidized LDLs activated NF- κ B, whereas long term incubation prevented NF- κ B activation triggered by pro-inflammatory agents.^[9] For instance, oxidized LDLs inhibit the LPS-induced NF- κ B activation and subsequent expression of TNF- α , IL-1 β and PAF-receptor in macrophages.^[9,64–67] In the same way, oxidized LDLs suppress the phytohaemagglutinin (PHA)-induced NF- κ B activation and subsequent IL-2 receptor expression in human T-lymphocytes.^[68]

Several bioactive lipids formed during LDL oxidation and present in oxidized LDLs could participate in this inhibitory effect, either by down-regulating signaling pathways implicated in the pro-inflammatory response or by activating antagonistic pathways.

Lyso-phosphatidylcholine (LPC) also regulates NF- κ B activity in a biphasic manner dependent on incubation time and concentration (NF- κ B being activated by low concentrations, but inhibited by high concentrations),^[37] similar to that observed with oxidized LDLs.^[9] The inhibitory effect of LPC was associated with a decreased expression of pro-inflammatory and pro-thrombotic proteins.^[69]

Oxidized phospholipids have been shown to reduce the endotoxin-induced tissue damages and to down-regulate LPS-induced cyclooxygenase-2 expression in human macrophages, possibly by targeting both ERK and NF- κ B pathways.^[67,70]

4-Hydroxy-2-nonenal (HNE), aldehydic molecules generated during peroxidation of PUFA, is also able to inhibit NF- κ B activation. For instance, HNE prevents the LPS-induced I κ B α phosphorylation in THP-1 monocytic cells,^[49] probably by reacting with and inhibiting IKK.^[71] HNE (like oxidized LDLs) is also able to inhibit the proteasome activity and subsequent proteolysis of I κ B α , possibly through a modification of proteasome proteins by 4-HNE.^[72,73] Finally, both mechanisms may converge to inhibit NF- κ B activation and subsequent expression of adhesion molecules, such as ICAM-1, VCAM-1 and E-selectin, in human aortic endothelial cells.^[74]

Oxidized PUFAs, such as 9-HODE and 13-HODE, prostaglandin J2 and HETE, have been identified as endogenous ligands and activators of PPAR γ that are able to reduce NF- κ B activation.^[75–77] PPAR α and PPAR γ are expressed in atherosclerotic lesions and can reduce adhesion molecules expression and recruitment of leukocytes to endothelial cells, regulate lipid metabolism in macrophagic cells, inhibit the proliferation and migration of smooth muscle cells and modulate the inflammatory response.^[77,78] PPARs antagonize NF- κ B activation by several converging mechanisms, including negative regulation of NF- κ B, overexpression of I κ B α and of antioxidant enzymes such as catalase.^[76,79,80]

Oxysterols have been shown to inhibit NF- κ B activation elicited by pro-inflammatory agents.^[65] This inhibitory effect could be mediated by transcription factors of the LXR and Sp families. LXRs are oxysterol-activated nuclear receptors that regulate genes controlling lipid homeostasis and inflammation.^[81,82] The anti-inflammatory effect of activated LXR is mediated, at least in part, through interference with the NF- κ B signaling pathway.^[83] Oxysterols are also able to regulate the activity of members of the Sp-family of transcription factors^[84] that are important regulators

of inflammatory processes.^[85] This family, belonging to the zinc-finger family of DNA binding proteins, consists of four proteins designed Sp1 (specific protein 1), Sp2, Sp3 and Sp4 which are general transcription factors that recognise GC and/or GT box motifs in the promoter of many ubiquitous or tissue specific genes.^[85] *In vitro* studies of the TNF- α promoter suggested that Sp1 and Sp3 may either compete with NF- κ B for binding to the promoter or interact with DNA-bound NF- κ B.^[84]

Potential Consequences of Oxidized LDLs-induced NF- κ B Modulation

NF- κ B plays a central role in host defense and inflammatory response, as well as in apoptosis regulation.^[86] Oxidized LDLs are able to alter the regulation, thereby altering the pro-/anti-inflammatory and pro-/anti-apoptotic balance in atherosclerotic plaques.^[87]

NF- κ B in Apoptosis and Survival

The analysis of RelA $-/-$, IKK β $-/-$ and IKK γ $-/-$ mice and cell biology studies have clearly shown that NF- κ B is an anti-apoptotic transcription factor.^[86] This also has been clearly demonstrated in the case of oxidized LDLs, using transfection of synthetic double-stranded DNA that binds and inhibits the transcriptional factor NF- κ B and thereby potentiates the pro-apoptotic effect induced by oxidized LDLs.^[88] The anti-apoptotic activity of NF- κ B is dependent on induction of genes^[86] such as members of the Bcl-2 family (A1, Bcl-xL), cellular inhibitors of apoptosis (c-IAPs), FLICE inhibitory protein (cFLIP), TNRF-associated factors TRAF1 and TRAF2 and A20 protein.^[86] These proteins act as inhibitors of both the death-receptor pathway (TRAF, cFLIP) and the mitochondrial apoptotic pathway (Bcl-2 related proteins, IAPs).^[86] Of note, most of the apoptosis inducers, such as oxidized LDLs, cytokines, TNF- α and chemotherapeutic agents, activate NF- κ B which in turn modulates the pro-apoptotic signaling.^[86,89,90] Reversely, during apoptosis, activated caspases cleave several anti-apoptotic proteins including NF- κ B itself, the NF- κ B regulatory proteins IKK, I κ B α (converted into a super-repressor-like molecule) and NF- κ B-regulated anti-apoptotic factors. This caspase-mediated proteolysis tends to eliminate the protection conveyed by NF- κ B survival pathway.^[86] In conclusion, NF- κ B is therefore at the crossroads of life and death and plays a critical role in it by regulating the level of anti-apoptotic proteins.^[86]

Although little evidence exists for a pro-apoptotic function of NF- κ B,^[86] in some specific cases, NF- κ B could induce the overexpression of various pro-apoptotic genes. For instance, in endothelial cells,

NF- κ B may enhance the pro-apoptotic effect of oxidized LDLs by up-regulating LOX-1 expression (LOX-1 promoter containing a consensus NF- κ B binding site),^[91,92] thereby increasing the uptake of oxidized LDLs and the subsequent toxicity. Another example is the oxidized LDL-induced NF- κ B-mediated overexpression of Angiotensin Receptor-1 (AT1) which may up-regulate LOX-1 expression and enhances the toxicity of oxidized LDLs^[93] and alter the balance between cell survival/apoptosis.^[94,95] Finally, NF- κ B may also up-regulate the expression of pro-apoptotic genes such as Fas ligand, Fas (Cd95), c-myc and p53, thereby sensitizing cells to apoptotic stimuli.^[96-98]

Adhesion Molecules and Cytokines

During atherogenesis, dyslipidemia and lipid deposition in the intima are coupled to a local inflammatory response, involving endothelial cells, monocytes and lymphocytes.^[99,100] Oxidized LDLs and other pro-inflammatory stimuli trigger a NF- κ B-mediated overexpression of adhesion molecules (VCAM-1, ICAM-1, E-selectin) that are involved in the "rolling" and adhesion of leukocytes to endothelial cells.^[101] Moreover, oxidized LDL also induced the expression of other pro-inflammatory genes of chemoattractant chemokines (MCP-1), growth factors (M-CSF, GM-CSF) and inflammatory cytokines (IL-1, TNF- α).^[102] As shown in Table I, these proteins of inflammation are, in part, under the control of NF- κ B.

Nitric Oxide (NO) Synthase and Endothelial Dysfunction

NO exhibits anti-inflammatory properties which is associated with inhibition of the expression of adhesion molecule (VCAM-1, ICAM-1, E-selectin) and cytokines (IL-8 and IL-6).^[100,103,104] This is mediated (at least in part) through inhibition (by NO) of NF- κ B activation, resulting either from overexpression and stabilization of the NF- κ B inhibitor I κ B α [105] and/or from a decreased DNA binding activity of NF- κ B dimers subsequent to the modification of the redox-sensitive Cys62 residue of p50 subunit.^[106] It may be noted that NO production (and subsequent anti-inflammatory and anti-atherogenic effects) are inhibited by oxidized LDLs and oxidized lipids.^[107-109]

Tissue Factor

Tissue factor (TF) is a potent pro-thrombotic protein that acts as a cofactor for activated factor VII (FVIIa) and induces thereby thrombin generation.^[110,111] It is expressed in vascular cells (monocytes, VSMCs, endothelial cells) in response to inflammatory cytokines, mitogens, endotoxins and initiates

rapidly the coagulation process in case of vascular injury.^[111,112] TF gene expression is regulated by a number of transcription factors that bind two distinct enhancers in the TF promoter. A distal enhancer, containing two AP-1 sites and one κB site controls TF expression mediated by LPS or cytokines, whereas a proximal enhancer, containing three overlapping Egr-1/Sp1 binding sites, controls TF expression induced by serum or phorbol ester. The concerted action of these different transcription factors allow to express TF in a cell- and stimulus-specific manner.^[112] It may be noted that oxidized LDLs and lipids can induce a biphasic effect on NF-κB activation and, subsequently, on TF level. This dual effect could explain apparently conflicting data reported in the literature. For instance, Brand *et al.*^[113] demonstrated that oxidized LDLs significantly enhanced TF expression induced by LPS in human monocytes, whereas other reports have shown that lysoPC induce a decrease in both NF-κB activation and TF expression induced by LPS.^[69]

Increased expression of TF is observed in all stages of atherosclerotic lesions, and plaques expressing the highest levels of TF are associated with the highest thrombotic risk.^[114] As oxidized lipids up-regulate TF expression in vascular cells^[115-118] and inactivate the anti-coagulant function of TFIP (tissue factor inhibitor protein),^[119,120] this results in an imbalance between TF and TFIP activities and, subsequently, in an increased thrombogenicity.

CONCLUSION

The biological responses induced by oxidized LDLs and mediated through NF-κB-regulated genes, are sometimes apparently puzzling, because of the dual effects of oxidized LDLs, summarized in Table II. This dual effect of oxidized LDLs may essentially result from two mechanisms: (i) modulation of

the balance between activation and inactivation of NF-κB (dependent on concentration of oxidized lipids and time of contact with the cells, inflammatory status of cells); (ii) stimulation by activated NF-κB of both the anti- and pro-apoptotic genes.

Activated NF-κB mediates, at least in part, the proinflammatory effects of oxidized LDLs, leading to the overexpression of adhesion molecules and chemokines, and subsequent recruitment and retention of mononuclear cells in the intima. This effect may be of particular importance in the early steps of atherogenesis and in the progression of atherosclerotic lesions.^[2-5,27,99,100] NF-κB activation also participates in the regulation of cell survival, and in the balance between proliferation and apoptosis induced by oxidized LDLs. On one hand, activation of NF-κB by oxidized LDLs activates anti-apoptotic genes that counterbalance the toxicity of oxidized LDLs, but, on the other hand, oxidized LDLs also induce a NF-κB-mediated up-regulation of LOX-1 and AT-1 receptors expression, which may enhance the pro-apoptotic effect of oxidized LDLs and angiotensin, respectively. Interestingly, both chronic and acute inflammatory sites co-exist with localized apoptosis at sites of plaque rupture, thus suggesting that the anti-apoptotic effect of NF-κB is unable to counterbalance the pathogenic effect of local inflammatory conditions in which NF-κB is implicated.^[27,78,100]

In contrast, oxidized LDLs are also able to reduce NF-κB activation induced by pro-inflammatory agents including LPS or cytokines. This inhibitory effect of oxidized LDLs may reduce the recruitment of leukocytes in the vascular wall, minimize the inflammatory response of activated T-lymphocytes and monocytes, impair the immune response against infections and limit TF expression. Down-regulation of the local immune responses can finally limit the inflammatory burst and maintain low level of inflammation which is a characteristic of the atherosclerotic lesions. But, on the other hand,

TABLE II Potential pathophysiological role of NF-κB in atherosclerotic lesions

(A) Potential consequences of NF-κB activation by oxLDL

Targets of NF-κB	Potential physiological consequences
Expression of adhesion molecules, chemotactic proteins, inflammatory cytokines	Inflammation
Up-regulation of Lox-1 and AT-II receptors	Apoptosis of endothelial cells Endothelial dysfunction
Survival pathway	Cell survival Plaque stability

(B) Potential consequences of NF-κB inhibition by oxLDL

Targets of NF-κB	Potential patho-physiological consequences
Reduced expression of inflammatory molecules (adhesion molecules, cytokines)	Immunosuppressive effect
Inhibition of Tissue Factor	Reduction of the pro-thrombotic state
Inhibition of iNOS expression and NO production	Pro- or anti-atherogenic effect depending on the local inflammatory status

NF- κ B inhibition reduces the expression of anti-apoptotic proteins, thereby sensitizing cells to toxic stimuli and increasing the instability of the plaque.

During the progression of atherosclerotic plaque, NF- κ B activation and the subsequent expression of anti-apoptotic proteins may favor healing and plaque stability, but may also lead to excessive fibro-proliferative process and luminal narrowing. Reversely, the inhibition of NF- κ B and subsequent anti-apoptotic proteins may elicit plaque instability and increase the risk of plaque erosion/rupture and subsequent athero-thrombosis.

The definition of the overall conditions that actually favour either the pro-atherogenic or the anti-atherogenic side of NF- κ B needs further investigation.

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